

Table 1

	Known SAPs		MS/MS from AdML S'some					MS/MS from Ftz S'some					
Reference with link	Protein Name	Mass	Avg XCorr	Total peptides	Unique sequences	Filtered peptides	Filtered unique	Avg XCorr	Total peptides	Unique sequences	Filtered peptides	Filtered unique	NCI annotation/GenBank Accession/Other Names

Sm core proteins

Sm core proteins													
SW:SMD1_HUMAN	Sm D1	13282	4.081	114	9	83	4	3.977	8	3	6	3	P13641
SW:SMD2_HUMAN	Sm D2	13527	3.897	50	14	26	5	4.012	15	8	10	5	P43330
SW:SMD3_HUMAN	Sm D3	13916	3.859	19	7	10	2	4.169	11	5	4	2	P43331
SW:RUXG_HUMAN	Sm G	8496	2.865	7	2	7	2	3.171	1	1	1	1	Q15357
SW:RSMB_HUMAN	Sm B/B'	24610	3.222	16	9	4	3	3.288	6	5	4	2	P14678
SW:RUXE_HUMAN	Sm E	10804	3.204	5	4	4	3	2.578	4	3	3	2	P08578
SW:RUXF_HUMAN	Sm F	9725	2.799	25	13	3	2	2.731	16	6	4	3	Q15356

U6 snRNA-associated sm-like proteins

UG snRNA-associated sm-like proteins													
SWN:LSM4_HUMAN	lsm4	15350	2.677	11	3	7	2	2.759	7	2	4	2	Q9y4z0
SWN:LSM2_HUMAN	lsm2	10835	4.074	9	6	4	2	3.435	3	2	3	2	Q9y333
SWN:LSM7_HUMAN	lsm7	11602	2.498	3	2	3	2	***					Q9uk45
SWN:LSM6_HUMAN	lsm6	9128	2.989	4	4	1	1	2.338	2	1	2	1	Q9y4y8
SWN:LSM3_HUMAN	lsm3	11714	2.24	10	6	1	1	2.457	6	2	3	1	Q9y4z1
SWN:LSM5_HUMAN	lsm5	9806	***					***					Q9Y4Y9
SWN:LSM8_HUMAN	lsm8	10271	***					***					Q95777

III. snRNP specific proteins

U1 snRNP specific proteins													
SW:RU17_HUMAN	U1 70k	70081	3.291	14	6	6	2	3.246	3	3	1	1	P08621
SW:RU1C_HUMAN	U1 C	17394	3.771	2	1	2	1	3.853	2	2	1	1	P09234
SW:RU1A_HUMAN	U1 A	31280	2.208	7	5	1	1	2.775	7	5	2	2	P09012

U2 snRNP specific proteins

GP:AF054284_1	SAP155	145915	3.475	124	43	64	23	2.743	52	26	29	15	
SW:S145_HUMAN	SAP145	97657	3.228	67	33	30	12	3.084	33	20	15	8	Q13435
SW:S114_HUMAN	SAP_114	88886	3.549	57	30	29	14	3.042	37	22	19	11	Q15459
GP:AJ001443_1	SAP_130	135592	3.471	16	10	9	5	3.096	9	6	7	5	
PIR2:A55749	SAP61	58849	3.308	33	21	20	11	2.921	15	7	7	3	
SW:SP62_HUMAN	SAP_62	49196	2.992	12	7	8	4	2.24	7	6	1	1	Q15428
SW:SP49_HUMAN	SAP49	44386	3.543	7	4	4	2	4.207	5	5	2	2	Q15427
SW:RU2A_HUMAN	U2 A'	28444	3.25	39	19	19	9	2.863	18	11	12	8	P09661
SW:RU2B_HUMAN	U2 B"	25486	4.024	11	6	6	1	5.28	6	3	1	1	
AAK94041	d14	14584	***					***					P08579

U5 snRNP specific proteins

G2 shRNA-specific proteins														
SWN:U520_HUMAN	U5-200	194479	3.395	150	62	101	36	2.929	154	56	96	32	lost link, DEIH-box	
GP-AB007510	U5-220	237785	0.337	111	42	59	25	2.778	143	53	77	29	lost link	
GNP-BC002360	U5-116	109478	3.308	70	38	47	22	2.812	68	34	45	16	EF-2-like	
GNP-BC001666	U5-102	106925	0.046	6	5	4	4	2.744	9	6	3	3	Prp6, hPrp1, SPF107	
GNP-BC002366	U5-100	95583	3.783	27	21	17	12	2.474	22	14	13	10	DEAD-box	
GP-AF090988	U5-40	39299	3.865	4	3	3	2	3.169	6	3	5	3		
SW:DM1_HUMAN	U5-15	16786	3.649	3	3	1	1	2.247	1	1	1	1	O14834, dm1 homolog	

U4/U6.U5 snRNP specific proteins

PIR2:T50839	U4/U6-90k	77529	3.335	15	13	9	8	2.558	11	8	6	5	hPrp3
GPN:BC007424_1	U4/U6-60k	58321	3.037	10	8	6	5	2.852	10	7	6	4	hPrp4
GP:AF083385_1	SPF30	26711	3.46	13	7	7	3	2.836	5	3	4	2	
PIR2:T00034	Tr-snRNP 110k	90255	3.147	22	16	11	11	2.918	19	15	11	9	SPF90/SART1/hSnu66
GP:AF353989_1	Tr-snRNP 65k	65415	2.46	2	2	1	1	***					hSad1p
SW:NHPX_HUMAN	Tr-snRNP15 5kD	14174	2.926	7	4	5	3	2.861	2	2	1	1	P55769, AF155235, hSnu13p
PIR1:S64705	CyP-60	58823	3.089	12	9	11	8	3.194	1	1	1	1	cyclophilin-like protein CyP-60
GPN:AF271652_1	PPIL3b	18155	3.162	9	6	5	4	3.185	5	5	2	2	cyclophilin-like protein PPIL3b (PPIL3)

Step II proteins

Step 1: proteins													
SW:PR16_HUMAN	hPrp16	140473	3.054	7	6	6	5	2.569	5	3	3	2	Q92620, DEAH-box
GP-AF038392	1 hPrp17	65521	3.2	10	6	6	4	2.585	3	2	1	1	
GPN-BC010634	1 hSlu7	68343	3.448	6	6	2	2	2.636	4	4	2	2	
GPN-BC000794	1 hPrp18	39860	4.918	1	1	1	1	***					
SW:DDX8_HUMAN	hPrp22	139315	3.364	12	9	10	7	2.71	15	13	9	8	Q14562, HRH1, DEAH-box

mRNP proteins

SWN:RBM8_HUMAN	Y14	19889	2.842	7	5	5	3	2.461	5	4	3	2	Q9y5s9, RBM8
AF047002	Aly	26861	3.32	8	6	2	1	2.841	9	6	3	2	NM_005782, BEF
SW:MGN_HUMAN	magoh	17164	2.934	7	5	3	2	2.713	7	4	5	3	P50606
PIR2:JC4525	RNPS1	34208	4.132	7	5	3	2	3.191	4	3	3	2	RNA-binding protein E5 1

SR proteins

SW:SFR1_HUMAN	SF2/ASF	27613	2.828	27	14	19	10	2.852	42	12	36	9	Q07955
SW:SFR3_HUMAN	SRp20	19330	2.762	16	4	11	3	2.764	11	4	8	3	P23152
SW:SFR7_HUMAN	9G8	27367	2.529	11	4	7	3	2.416	6	3	4	3	Q16629
SW:SFR6_HUMAN	SRp55	39568	3.181	6	4	4	3	3.027	8	4	6	3	Q13247
SW:SFR4_HUMAN	SRp75	56792	2.381	5	4	3	2	2.537	6	3	2	2	Q08170
SW:SFR5_HUMAN	SRp40	31264	3.327	2	2	2	2	3.018	4	1	4	1	Q13243
SW:SFR9_HUMAN	SRp30c	25542	2.759	5	3	2	2	2.414	10	6	5	4	Q13242
SW:SFR2_HUMAN	SC35	25575	2.974	2	2	1	1	***					Q01130
GP:AF048977_1	SRm160	93519	3.559	5	4	2	2	3.578	7	5	5	3	
SW:SFRB_HUMAN	p54	53542	3.633	9	5	5	3	2.276	2	2	1	1	Q05519, SFRS11
SW:SFR8_HUMAN	SWAP	104821	***					2.282	4	4	1	1	Q12872, SFRS8

Other dead box/helicase proteins

SW:HE47_HUMAN	UAP56	48991	3.131	7	7	3	3	2.54	7	5	5	3	hBAT1, p47, HE47, Q13838
SW:DD15_HUMAN	hPrp43	92829	3.237	14	11	12	9	2.778	11	10	5	4	Q43143
SW:DD17_HUMAN		72371	2.894	13	10	11	8	2.554	20	11	14	6	Q92841, p72
SW:DDX3_HUMAN		73243	3.233	6	5	6	5	2.54	3	3	2	2	Q00571, nlp2
SW:DDX5_HUMAN		69148	2.643	10	7	6	6	2.515	9	6	5	3	P17844, p68, mann
SW:DD16_HUMAN	hPrp2	119172	2.64	10	7	6	5	2.613	9	6	5	3	Q60231
SW:DDX9_HUMAN		140877	3.063	7	6	2	1	2.75	9	7	6	5	Q08211, ndh ii
GP:AF106680_1	hPrp5	117266	2.539	2	2	1	1	3.455	1	1	1	1	RNA helicase

Other SAPs

SW:CB80_HUMAN	CBP 80	18139	2.846	28	16	20	12	2.672	38	18	20	10	Q09161
SW:CB20_HUMAN	CBP20	18001	3.281	8	4	2	1	2.482	8	5	4	3	P52298
SW:U2AF_HUMAN	U2 AF65	53501	3.641	15	8	11	5	3.758	6	5	1	1	P26368
SW:U2AG_HUMAN	U2AF35	27872	2.77	3	3	2	2	3.942	2	2	2	2	Q01081
GN:AY029347_1	hPrp4/kinase	116973	3.838	6	3	5	2	2.967	3	3	2	2	serine/threonine-protein kinase (PRP4)
GN:BC008719_1	hPrp19	55181	3.165	43	17	19	9	2.931	20	10	12	7	nuclear matrix protein NMP200, WD repeat
GP:AL050389_1	hPrp31	55424	3.049	8	6	4	3	3.207	4	4	3	3	
GP:AF049523_1	hFBP 11		2.917	8	4	7	3	2.864	11	5	6	3	huntingtin-interacting protein HYPA/EBP11
GP:AF255443_1	hCRN	99201	3.032	25	16	16	11	3.318	8	6	4	4	CGI-201, TPR-repeat
Y08765/Y08766	SFI1/mBBP	68632	***					***					several isoforms

SAPs with other function

PIR2:T08599	CA150	123960	2.925	56	28	33	15	2.684	32	23	21	13	
SW:SKIP_HUMAN	skip	61494	3.507	22	18	8	7	2.896	18	9	12	5	snw1, nuclear receptor coactivator ncoa-62
GN:BC007871_1	SPF45	44962	2	2	1	1		3.118	1	1	1	1	G-patch, RRM, DNA damage repair
GP:AF083383_1	SPF38	34290	2.876	4	3	2	2	3.823	5	4	4	3	WD

Proteins enriched in H complex

SW:PTB_HUMAN	hnRNP I/PTB	57221	4.13	25	18	11	6	2.206	3	3	1	1	P26599
SW:ROA1_HUMAN	hnRNP A1	38715	3.48	15	11	8	6	3.354	20	13	8	5	P09651
SW:ROA2_HUMAN	hnRNP A2/B1	37430	3.515	3	3	2	2	2.653	14	8	8	4	P22626
SWN:ROR_HUMAN	hnRNP R	70943	3.074	10	9	7	7	3.006	15	9	11	6	Q43390
SW:ROC_HUMAN	hnRNP C1/C2	33299	2.828	7	6	6	6	2.791	34	10	27	10	P07910
SW:ROK_HUMAN	hnRNP K	50976	3.482	5	5	5	5	3.356	16	9	9	6	Q07244
SW:ROA0_HUMAN	hnRNP A0	30841	4.284	7	3	4	2	4.373	3	2	3	2	Q13151
SW:ROH1_HUMAN	hnRNP H	49229	3.47	6	5	4	3	***					P31943
SW:ROL_HUMAN	hnRNP L	60187	2.884	6	6	3	3	2.989	55	21	37	14	P14866
SWN:ROD_HUMAN	hnRNP D0	38434	2.879	4	3	3	2	3.365	8	5	4	1	Q14103
SW:ROM_HUMAN	hnRNP M	77469	2.815	5	5	3	3	***					P52272
GN:BC001616_1	hnRNP A/B	30588	3.667	2	1	2	1	3.06	1	1	1	1	
SW:ROA3_HUMAN	hnRNP A3	39686	3.655	6	5	2	2	2.823	7	6	3	2	P51991
SW:ROF_HUMAN	hnRNP F	45672	3.429	2	2	2	2	***					P52597
SW:ROG_HUMAN	hnRNP G	42404	***					3.036	10	4	7	3	P38159
PIR2:B54857	NF-AT 90k	73339	3.117	7	4	7	4	2.919	9	8	7	6	
PIR2:A54857	NF-AT 45k	44697	3.436	7	6	4	4	3.197	8	7	6	5	
GP:AF037448_1	Gry-rbp	69633	3.882	3	3	1	1	3.149	9	5	6	4	RRM RNA binding protein
GN:BC008875_1	PUF60	58171	3.316	57	29	24	13	3.217	29	18	5	4	siah binding protein 1, FBP interacting repressor, PTB

Table 2

	Putative Novel SAPs		MS/MS from AdML- S'some					MS/MS from Fit S'some				
Reference with link	NCI annotation	Mass	Avg XCorr	Total peptides	Unique sequences	Filtered peptides	Filtered unique	Avg XCorr	Total peptides	Unique sequences	Filtered peptides	Filtered unique
Potentially new SAPs												
GP:AF356524_1	nuclear receptor transcription cofactor (SHARP)	402248	3.39	32	25	16	13	2.829	23	18	10	8
GPN:BC007208_1	HCNP, XPA-binding protein 2	100010	3.241	27	18	17	12	2.779	9	8	6	5
GP:AC004858_3	U1 small nonnucleoprotein 1SNRP homolog, Pole(A) binding protein	94122	3.331	50	26	33	15	3.454	24	15	13	8
PIR2:155595	splicing factor	58657	3.346	32	14	18	7	2.9	10	5	4	2
GP:AB034205_1	osipatin resistance-associated overexpressed protein	51466	3.45	14	9	10	6	3.149	8	7	7	6
SW:BUB3_HUMAN	mitotic checkpoint protein bub3	37155	2.982	6	5	6	5	2.956	7	4	5	3
SWN:ELV1_HUMAN	elav-like protein 1 (hu-antigen r) (hur)	36062	2.787	2	2	1	1	2.93	10	6	5	3
GPN:BC001621_1	Npw38-binding protein NpwBP	69998	2.64	11	9	7	5	2.332	5	4	2	1
SW:RED_HUMAN	red protein (rer protein) (ik factor) (cytokine ik)	65630	2.781	9	7	5	4	2.517	4	4	2	2
SW:Z07_HUMAN	O43670, zinc finger protein 207	50751	3.199	10	6	2	2	2.64	6	5	1	1
SW:YB1_HUMAN	y box binding protein-1 (yb-1) (ccaaat-binding transcription factor 1 subunit a) (cbf-a) enhancer factor	35924	4.636	10	9	1	1	4.316	6	5	2	1
GPN:BC003376_1	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)	36092	3.878	2	1	1	1	3.862	5	1	3	1
PIR2:A53545	nuclear matrix protein p84	75627	3.557	13	10	6	5	2.99	29	13	15	7
GP:AF155096_1	NY-REN-6 antigen, partial cds		2.973	17	6	4	2	2.82	11	4	6	3
SW:IF4N_HUMAN	eukaryotic initiation factor 4a-like nuk-34	46833	3.147	14	12	9	9	2.383	11	8	6	5
SWN:CRK7_HUMAN	cell division cycle 2-related protein kinase 7 (cdc2-related protein kinase 7) (crkrs)	164155	3.146	11	8	6	3	3.07	7	6	5	4
GPN:BC001403_1	pre-mRNA cleavage factor Im (25kD)	26227	3.246	11	7	7	5	3.511	6	5	4	3
GP:AF044333_1	pleiotropic regulator 1 (PLRG1)	57194	3.768	13	10	7	6	3.656	4	4	3	3
SW:G10_HUMAN	P41223, human homolog of xenopus maternal g10 protein (edg-2)	16844	2.277	2	2	1	1	2.431	4	2	3	1
GP:AJ271745_1	double-stranded RNA binding nuclear protein DRBP76 (ILF3 gene)	76033	2.804	2	1	1	1	2.41	5	3	3	1
GP:AJ276706_1	partial mRNA for WTAP (wilm's tumor associating protein) (hFL2D)		3.572	2	1	1	1	3.715	2	2	2	2
GP:AJ279080_1	putative transcription factor (ORF1), ORF1 S1 RNA binding protein	104804	3.544	3	3	3	3	3.033	3	3	2	2
SW:GCFC_HUMAN	gc-rich sequence dna-binding factor homolog, Q95b6	29010	3.167	3	3	3	3	2.655	2	1	2	1
GP:U70667_1	Fas-ligand associated factor 1 (FLAF1), WWP/VW motif		2.51	10	6	3	1	2.616	5	4	2	2
GPN:BC005152_1	similar to mouse Gli3 or D. melanogaster transcription factor IIB	22774	3.154	1	1	1	1	2.606	2	2	2	2
SWN:R856_HUMAN	Q92804, tata-binding protein associated factor 2n (ma-binding protein 56) (tbf68)	61830	2.726	3	2	3	2	2.011	3	2	2	1
SWN:CYCK_HUMAN	O75909, cyclin k	41293	3.653	2	1	1	1	4.355	2	2	1	1
GPN:BC003015_1	Similar to expressed sequence 2 embryonic lethal		3.363	3	3	3	3	4.199	1	1	1	1
GP:AB016088_1	RNA binding protein, partial cds		3.764	10	5	5	2	4.054	3	3	1	1
SW:SP18_HUMAN	O00422, sin3 associated polypeptide p18	17561	3.414	2	2	1	1	3.435	5	5	1	1
GPN:BC002548_1	Similar to Moloney leukemia virus 10	113671	3.215	4	4	2	2	3.434	2	2	1	1
PIR2:138191	nucleic acid binding protein (fragment)		4.65	1	1	1	1	3.364	2	2	1	1
SW:CIRP_HUMAN	cold-inducible ma-binding protein (glycine-rich ma-binding protein grp)	18648	3.037	5	3	4	2	3.349	2	2	1	1
GP:AF112222_1	nuclear protein SDK3	81584	3.026	4	3	1	1	3.311	5	3	1	1
GP:L76159_1	Facioscapulohumeral muscular dystrophy region gene-1	29172	3.363	2	1	2	1	3.079	1	1	1	1
SW:GR78_HUMAN	78 kda glucose-regulated protein precursor (grp 78) (immunoglobulin heavy chain binding protein) (bip)	72116	4.29	3	2	1	1	2.955	1	1	1	1
GPN:BC000495_1	CD2 antigen (cytoplasmic tail)-binding protein 2	37646	3.374	4	3	2	2	2.887	4	4	1	1
SW:MFA1_HUMAN	P55081, microfilament-associated protein 1	51855	2.687	6	5	3	3	2.777	3	3	1	1
GP:AF015044_1	EH-binding protein, binds EH domains of eps15		3.132	6	6	3	3	2.771	1	1	1	1
GPN:BC010381_1	nuclear matrix protein p84	75666	2.555	5	5	3	3	2.768	6	4	1	1
GP:AF361746_1	endothelial cell-selective adhesion molecule (ESAM), immunoglobulin superfamily, contains V and C2 domains	41208	3.24	3	1	3	1	2.555	1	1	1	1
GP:AJ271741_1	partial ILF3 gene for interleukin enhancer binding factor 3	25972	2.303	3	2	2	1	2.379	4	3	1	1
SW:EF11_HUMAN	elongation factor 1-alpha 1 (ef-1-alpha-1) (elongation factor tu) (ef-tu)	50141	2.852	6	5	5	4	2.377	1	1	1	1
SW:DBPA_HUMAN	dna-binding protein a (cold shock domain protein a) (single-strand dna binding protein nf- gmb)	40060	3.194	10	8	5	3	2.374	2	2	1	1
Hypothetical new SAPs												
PIR2:T00365_??	hypothetical protein KIAA0670 (fragment)		3.207	77	37	44	18	3.001	72	33	45	17
PIR2:T02345	hypothetical protein KIAA0324 (fragment), SRm300, matrix related		3.069	58	30	31	17	2.885	37	25	20	13
SW:Y017_HUMAN	hypothetical protein kaa0017	44806	3.668	39	20	24	9	3.192	19	11	13	5
GP:AB007892_1	KIAA0432 mRNA	58171	3.417	51	32	23	16	2.962	27	21	15	12
PIR2:T00333	hypothetical protein KIAA0560	163986	3.292	31	21	18	12	2.643	30	19	17	12
PIR2:T12455	hypothetical protein DKFZp564H2023.1 (fragments)		2.898	35	18	20	10	2.931	67	21	40	10
SW:YS64_HUMAN	hypothetical protein s164 (fragment)		3.614	12	8	7	4	2.998	16	7	12	6

GP:AK023659_1	clone PLACE1009798, weakly similar to RLR1 PROTEIN	73795	2.902	12	8	11	7	2.698	33	16	25	12
PIR2:T46386	hypothetical protein DKFZp434P011.1 (fragment)		3.071	15	11	11	8	2.602	17	12	12	9
GN:BC003118_1	clone MGC:2655 IMAGE 3537243, mRNA, complete cds	34849	3.259	5	3	4	2	2.565	14	6	11	4
GP:AF151059_1	HSPC225 (FBP11 related)	24297	2.938	14	6	8	5	2.66	15	5	10	4
PIR2:I39463	gene anonymous protein - human	78536	3.161	5	4	3	2	2.629	16	12	8	6
SWN:CG80_HUMAN	hypothetical protein cgi-110 (protein hspc175)	14585	3.8	13	7	7	2	3.082	5	4	4	3
GN:BC006397_1	Similar to hypothetical protein FLJ12479, clone MGC:13150 IMAGE:4298786	102135	3.052	9	5	4	2	2.882	7	6	4	3
GP:AL031668_1	Human DNA sequence from clone RP1-64K7 on chromosome 20q11.21-11.23 Contains the EIF2S2 gene for eukaryotic translation initiation factor 2 subunit 2 (beta, 38kD), a putative novel gene, the gene for heterogenous nuclear ribonucleoprotein RALY or aut	32214	2.707	2	2	1	1	2.547	5	3	4	2
GP:D38552_1	KIAA0073 gene, partial cds, The ha1539 protein is related to cyclophilin		2.924	14	10	9	7	2.585	8	6	4	4
GP:AB046824_1	Homo sapiens mRNA for KIAA1604 protein, partial cds, Start codon is not identified		3.596	16	11	11	8	3.345	8	6	3	3
GP:AF130096_1	FLC0586 PRO2855 mRNA, complete cds	34655	3.323	10	4	8	3	3.174	3	2	3	2
GN:BC003048_1	CGI-124 protein	18237	3.038	8	4	7	4	2.716	4	3	3	2
GP:AB018344_1	Homo sapiens mRNA for KIAA0801 protein, complete cds	117461	3.37	24	14	14	9	2.586	10	9	3	2
GP:AK027098_1	Homo sapiens cDNA: FLJ23445 fis, clone HSI01721, unnamed protein product	23671	2.498	4	3	3	2	2.566	10	6	3	3
GN:BC003402_1	hypothetical protein FLJ10290, clone MGC:4943 IMAGE:3449258, mRNA, complete cds	46896	3.376	8	8	3	3	2.5	5	4	3	2
PIR2:T12485	hypothetical protein DKFZp564O2082.1 - human	28722	4.171	4	4	1	1	3.224	6	4	2	2
GN:BC004442_1	Similar to RIKEN cDNA 5830446M03 gene, clone MGC 4036 IMAGE:2820683, mRNA, complete	32992	2.902	4	4	3	3	3	2	1	2	1
PIR2:T17232	hypothetical protein DKFZp434I116.1 - human (fragment)		3.077	4	4	2	2	2.956	4	4	2	2
GN:BC006474_1	clone IMAGE 2820942, mRNA, partial cds		3.342	2	2	2	2	2.813	4	4	2	2
GN:BC004122_1	Similar to RIKEN cDNA 5830446M03 gene, clone MGC:11203 IMAGE 3927759, mRNA	10870	3.599	2	2	1	1	2.707	2	1	2	1
SW:Y105 HUMAN	hypothetical protein kaa0105	17801	3.049	3	3	3	3	2.563	2	1	2	1
GN:BC002876_1	hypothetical protein FLJ10806	57544	3.093	12	6	8	4	2.561	3	2	2	1
GP:AF161497_1	Homo sapiens HSPC148 mRNA, complete cds	26610	2.734	5	3	2	1	2.403	5	3	2	2
PIR2:T12531	hypothetical protein DKFZp434B194.1	96820	3.006	10	10	5	5	2.336	7	6	2	2
GP:AB023146_1	KIAA0929 protein, partial cds		3.205	22	18	15	11	2.015	9	6	2	2
GN:BC003359_1	hypothetical protein, clone MGC 5267 IMAGE 2900332, mRNA, complete cds	33855	2.687	2	2	1	1	4.451	1	1	1	1
GN:BC000198_1	Similar to CG11985 gene product, clone MGC:3133 IMAGE 3352960, mRNA, complete cds	10135	3.837	33	2	4	1	4.05	2	1	1	1
PIR2:T02672	hypothetical protein R31449_3 - human (fragment)		3.456	6	6	3	3	3.604	4	4	1	1
PIR2:T46935	hypothetical protein DKFZp434D199.1	14263	3.763	6	6	4	4	3.106	1	1	1	1
GP:AF161433_1	HSPC315 mRNA, partial cds		3.043	3	2	2	1	2.955	2	2	1	1
GP:AF132955_1	CGI-21 protein mRNA	37542	2.706	4	2	2	1	2.795	1	1	1	1
GP:AK000741_1	cDNA FLJ20734 fis, clone HEP08523, unnamed protein product	70516	3.219	3	3	1	1	2.692	2	2	1	1
SW:Y052 HUMAN	hypothetical protein kaa0052 (fragment)		2.916	6	6	3	3	2.611	3	3	1	1
GP:AB011132_1	KIAA0560 protein, partial cds		2.525	1	1	1	1	2.595	1	1	1	1
GN:BC004258_1	Homo sapiens, hypothetical protein PRO1741, clone MGC:10753 IMAGE:3347345, mRNA, complete cds	65691	3.813	3	2	2	2	2.567	1	1	1	1
GN:BC006350_1	clone MGC:13125 IMAGE:4111572	70521	2.77	9	9	7	7	2.419	1	1	1	1
GP:AL512685_1	cDNA DKFZp547K202 (from clone DKFZp547K202); WD-protein The frame shift was determined manually	75399	3.222	3	3	1	1	2.373	4	4	1	1
GP:AL023804_1	Human DNA sequence from clone RP4-633O20 on chromosome 20q11.23-12 Contains 5' end of a gene similar to Bos taurus P14 protein (P14L), ESTs, CA repeat(D20S859), STSs and GSSs, complete sequence, Also similar to Drosophila CG11964 protein	32895	2.81	4	4	3	3	2.321	2	2	1	1